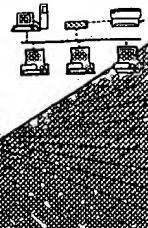


BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/922,549A
Source: OIPG
Date Processed by STIC: 2/5/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>) , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

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Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 09/922,549A
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 _____ Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 _____ Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 _____ Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4 _____ Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 _____ Variable Length	Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 _____ PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 _____ Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 _____ Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 _____ Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.	
10 _____ Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 ✓ _____ Use of <220>	Sequence(s) <u>32</u> missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 _____ PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 _____ Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.	

OIPE

RAW SEQUENCE LISTING

DATE: 02/05/2002

PATENT APPLICATION: US/09/922,549A

TIME: 11:44:05

Input Set : A:\Lysozyme.ST25.txt

Output Set: N:\CRF3\02052002\I922549A.raw

3 <110> APPLICANT: Avigenics
 5 <120> TITLE OF INVENTION: Chicken Lysozyme Promoter
 7 <130> FILE REFERENCE: A181 8060
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/922,549A
 C--> 9 <141> CURRENT FILING DATE: 2002-01-10
 9 <160> NUMBER OF SEQ ID NOS: 65
 11 <170> SOFTWARE: PatentIn version 3.0
 13 <210> SEQ ID NO: 1
 14 <211> LENGTH: 20
 15 <212> TYPE: DNA
 C--> 16 <213> ORGANISM: artificial
 18 <220> FEATURE:
 19 <223> OTHER INFORMATION: Primer 5pLMAR2
 21 <400> SEQUENCE: 1
 22 tgccgccttc ttgatattc
 25 <210> SEQ ID NO: 2
 26 <211> LENGTH: 20
 27 <212> TYPE: DNA
 C--> 28 <213> ORGANISM: artificial
 30 <220> FEATURE:
 31 <223> OTHER INFORMATION: Primer LE-6.1kbrev1
 33 <400> SEQUENCE: 2
 34 ttggtggttaa ggcctttttg
 37 <210> SEQ ID NO: 3
 38 <211> LENGTH: 20
 39 <212> TYPE: DNA
 C--> 40 <213> ORGANISM: artificial
 42 <220> FEATURE:
 43 <223> OTHER INFORMATION: Primer lys-6.1
 45 <400> SEQUENCE: 3
 46 ctggcaagct gtcaaaaaca
 49 <210> SEQ ID NO: 4
 50 <211> LENGTH: 20
 51 <212> TYPE: DNA
 C--> 52 <213> ORGANISM: artificial
 54 <220> FEATURE:
 55 <223> OTHER INFORMATION: Primer LysElrev
 57 <400> SEQUENCE: 4
 58 cagctcacat cgtccaaaga
 61 <210> SEQ ID NO: 5
 62 <211> LENGTH: 34
 63 <212> TYPE: DNA
 C--> 64 <213> ORGANISM: artificial

p6
 Does Not Comply
 Corrected Diskette Needed

20

20

20

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RAW SEQUENCE LISTING

DATE: 02/05/2002

PATENT APPLICATION: US/09/922,549A

TIME: 11:44:05

Input Set : A:\Lysozyme.ST25.txt

Output Set: N:\CRF3\02052002\I922549A.raw

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66 <220> FEATURE:
67 <223> OTHER INFORMATION: Primer LYSBSU
69 <400> SEQUENCE: 5
70 ccccccccta aggcagccag gggcaggaag caaa 34
73 <210> SEQ ID NO: 6
74 <211> LENGTH: 12
75 <212> TYPE: DNA
C--> 76 <213> ORGANISM: artificial
78 <220> FEATURE:
79 <223> OTHER INFORMATION: Primer SaltoNotI
81 <400> SEQUENCE: 6
82 tcgagcggcc gc 12
85 <210> SEQ ID NO: 7
86 <211> LENGTH: 20
87 <212> TYPE: DNA
C--> 88 <213> ORGANISM: artificial
90 <220> FEATURE:
91 <223> OTHER INFORMATION: Primer T7
93 <400> SEQUENCE: 7
94 taatacgact cactataggg 20
97 <210> SEQ ID NO: 8
98 <211> LENGTH: 21
99 <212> TYPE: DNA
C--> 100 <213> ORGANISM: artificial
102 <220> FEATURE:
103 <223> OTHER INFORMATION: Primerlys61enfor1
105 <400> SEQUENCE: 8
106 cgtggtgatc aaatctttgt g 21
109 <210> SEQ ID NO: 9
110 <211> LENGTH: 20
111 <212> TYPE: DNA
C--> 112 <213> ORGANISM: artificial
114 <220> FEATURE:
115 <223> OTHER INFORMATION: Primer lys61enrev1
117 <400> SEQUENCE: 9
118 aggagggcac agtagggatc 20
121 <210> SEQ ID NO: 10
122 <211> LENGTH: 19
123 <212> TYPE: DNA
C--> 124 <213> ORGANISM: artificial
126 <220> FEATURE:
127 <223> OTHER INFORMATION: Primer 5MARfor1
129 <400> SEQUENCE: 10
130 gtggcctgtg tctgtgctt 19
133 <210> SEQ ID NO: 11
134 <211> LENGTH: 20
135 <212> TYPE: DNA
C--> 136 <213> ORGANISM: artificial
138 <220> FEATURE:

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RAW SEQUENCE LISTING

DATE: 02/05/2002

PATENT APPLICATION: US/09/922,549A

TIME: 11:44:05

Input Set : A:\Lysozyme.ST25.txt

Output Set: N:\CRF3\02052002\I922549A.raw

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139 <223> OTHER INFORMATION: Primer IFN-3rev
141 <400> SEQUENCE: 11
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145 <210> SEQ ID NO: 12
146 <211> LENGTH: 20
147 <212> TYPE: DNA
C--> 148 <213> ORGANISM: artificial
150 <220> FEATURE:
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153 <400> SEQUENCE: 12
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157 <210> SEQ ID NO: 13
158 <211> LENGTH: 20
159 <212> TYPE: DNA
C--> 160 <213> ORGANISM: artificial
162 <220> FEATURE:
163 <223> OTHER INFORMATION: Primer lys002for
165 <400> SEQUENCE: 13
166 ctctcagaat gcccaactcc 20
169 <210> SEQ ID NO: 14
170 <211> LENGTH: 20
171 <212> TYPE: DNA
C--> 172 <213> ORGANISM: artificial
174 <220> FEATURE:
175 <223> OTHER INFORMATION: Primer lys003for
177 <400> SEQUENCE: 14
178 tgtattggtc tccctcctgc 20
181 <210> SEQ ID NO: 15
182 <211> LENGTH: 20
183 <212> TYPE: DNA
C--> 184 <213> ORGANISM: artificial
186 <220> FEATURE:
187 <223> OTHER INFORMATION: Primer lys005for
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190 tgttgaaatt gcagtgtggc 20
193 <210> SEQ ID NO: 16
194 <211> LENGTH: 20
195 <212> TYPE: DNA
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198 <220> FEATURE:
199 <223> OTHER INFORMATION: Primer lys006rev
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206 <211> LENGTH: 20
207 <212> TYPE: DNA
C--> 208 <213> ORGANISM: artificial
210 <220> FEATURE:
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RAW SEQUENCE LISTING

DATE: 02/05/2002

PATENT APPLICATION: US/09/922,549A

TIME: 11:44:05

Input Set : A:\Lysozyme.ST25.txt

Output Set: N:\CRF3\02052002\I922549A.raw

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218 <211> LENGTH: 20
219 <212> TYPE: DNA
C--> 220 <213> ORGANISM: artificial
222 <220> FEATURE:
223 <223> OTHER INFORMATION: Primer lys008rev
225 <400> SEQUENCE: 18
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229 <210> SEQ ID NO: 19
230 <211> LENGTH: 20
231 <212> TYPE: DNA
C--> 232 <213> ORGANISM: artificial
234 <220> FEATURE:
235 <223> OTHER INFORMATION: Primer lys009for
237 <400> SEQUENCE: 19
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242 <211> LENGTH: 20
243 <212> TYPE: DNA
C--> 244 <213> ORGANISM: artificial
246 <220> FEATURE:
247 <223> OTHER INFORMATION: Primer lys010rev
249 <400> SEQUENCE: 20
250 tccatggtgg tcaaacagaa                                20
253 <210> SEQ ID NO: 21
254 <211> LENGTH: 20
255 <212> TYPE: DNA
C--> 256 <213> ORGANISM: artificial
258 <220> FEATURE:
259 <223> OTHER INFORMATION: Primer lys011for
261 <400> SEQUENCE: 21
262 gtactagacc aggcagccca                                20
265 <210> SEQ ID NO: 22
266 <211> LENGTH: 20
267 <212> TYPE: DNA
C--> 268 <213> ORGANISM: artificial
270 <220> FEATURE:
271 <223> OTHER INFORMATION: Primer lys012rev
273 <400> SEQUENCE: 22
274 gtgggaagta ccacattggc                                20
277 <210> SEQ ID NO: 23
278 <211> LENGTH: 20
279 <212> TYPE: DNA
C--> 280 <213> ORGANISM: artificial
282 <220> FEATURE:
283 <223> OTHER INFORMATION: Primer lys013for
285 <400> SEQUENCE: 23

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RAW SEQUENCE LISTING

DATE: 02/05/2002

PATENT APPLICATION: US/09/922,549A

TIME: 11:44:05

Input Set : A:\Lysozyme.ST25.txt

Output Set: N:\CRF3\02052002\I922549A.raw

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286 cgctcaggag aaagtgaacc                                20
289 <210> SEQ ID NO: 24
290 <211> LENGTH: 20
291 <212> TYPE: DNA
C--> 292 <213> ORGANISM: artificial
294 <220> FEATURE:
295 <223> OTHER INFORMATION: Primer lys014rev
297 <400> SEQUENCE: 24
298 cggttttgcc tttgtgtttt                                20
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302 <211> LENGTH: 20
303 <212> TYPE: DNA
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306 <220> FEATURE:
307 <223> OTHER INFORMATION: Primer lys015rev
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314 <211> LENGTH: 20
315 <212> TYPE: DNA
C--> 316 <213> ORGANISM: artificial
318 <220> FEATURE:
319 <223> OTHER INFORMATION: Primer lys016rev
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322 gccaatcaga ctgcatttca                                20
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326 <211> LENGTH: 20
327 <212> TYPE: DNA
C--> 328 <213> ORGANISM: artificial
330 <220> FEATURE:
331 <223> OTHER INFORMATION: Primer lys017rev
333 <400> SEQUENCE: 27
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339 <212> TYPE: DNA
C--> 340 <213> ORGANISM: artificial
342 <220> FEATURE:
343 <223> OTHER INFORMATION: Primer lys018for
345 <400> SEQUENCE: 28
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349 <210> SEQ ID NO: 29
350 <211> LENGTH: 20
351 <212> TYPE: DNA
C--> 352 <213> ORGANISM: artificial
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355 <223> OTHER INFORMATION: Primer lys019rev
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<210> 32

<211> 20

<212> DNA

<213> Artificial

<400> 32

atttgctgtg gtggatgtga

20

see item 11 on Ena Summary Sheet

use Use of n and/or Xaa has been detected in the Sequence Listing.

Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/922,549A

DATE: 02/05/2002

TIME: 11:44:06

Input Set : A:\Lysozyme.ST25.txt

Output Set: N:\CRF3\02052002\I922549A.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
 L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
 L:16 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
 L:28 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
 L:40 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
 L:52 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
 L:64 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
 L:76 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
 L:88 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
 L:100 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
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 L:376 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:31
 L:388 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:32
 L:390 M:258 W: Mandatory Feature missing, <220> FEATURE:
 L:390 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
 L:397 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:33
 L:409 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:34
 L:421 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:35
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/922,549A

DATE: 02/05/2002

TIME: 11:44:06

Input Set : A:\Lysozyme.ST25.txt

Output Set: N:\CRF3\02052002\I922549A.raw

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L:577 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:48
L:589 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:49
L:601 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:50